FIGURE 1

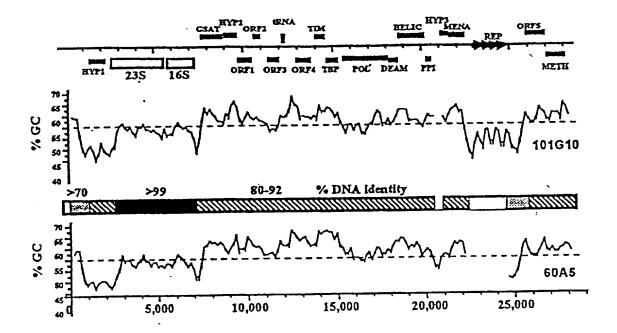


Figure 2

60 0. 00.	Gene	<u>Strain</u>		TATA Box			Coding Start		TATA to Start (bp)	
81 82	Hypoth 03	A B					GCGGCGCATG CCGGCGCGTG			25
83 84	Hypoth 02	A B					CGGGGCCCAT			26
85 96	ORF 02	A B					AACGGCCGTA ACCTGCCGTA			27
88	ORF 03	A B	CATGGAACTA	GATAAT	AACC	GGTCCCGCGG	ATCCCATGCA GTACAATGCA	T G	~~~~~~~	27
89 90 91	PPI	A B	AGCACGACAA	GTTATA	GCAG	GGTACAAAGG	GTGCGCGCGC AGCAGCGCAC	ATG~~~~~	~~~~~~~~	28
92	GSAT	A B	ATCCGGCCTC	ATTAAA	TTAC	GGGGGTACA	GCCTGCTGCC ACCTGCTGCC	GTG~~~~~	~~~~~~~	28
99	ORF 05	A B	ACTTCATACA	CATAAA	TCCC	GCCTGAACGG	GCGGCTGCGC TCGTCCGCGC	ATG~~~~~	~~~~~~~	28
96	deaminase	A B A	CCGCATATAC	CATAAT	ATGC	CGGGCGGGG	CACCATGGCC CAGGCTGCCC CAGGGCCGCG	.GTG~~~~~	~~~~~~~	29
98		B A	GGGTAGAAAC	CATAAA	ACAA	CAGGCCGCGG	CAGGGCCGCG CAGGGCG.CG GCGCGTATCA	CGTG~~~~~	~~~~~~~	29
100	tRNA-tyr	B A	ATACACGTGG	TATAAA	CAGA	GG.CCGGACG	GCGCGGACCA CACGGATCGT	CATG~~~~	~~~~~~~	29
103		B A	GCGATAGTTA	TTTAAA	ACTA	GGATGCCGGG	CACCCGTCGT GGATCCTGAC	CCCA~~~~	~~~~~~~	30
105	TIM	B A	CCGGGCCCCG GCGTCGATAG	GTTAAA AATAAA	ATAG TACG	AGTGCGGCCG CGCAGGGGGC	GGCACCGGAT CCCGTGGCGC	CAATG~~~~ GATCGCCCGT		36
107	Hypoth 01	B A	ATTTCAACTA	CATAAA	TGCC	TAGTTACGCA	GCGGTGC GAAATAGCAA	ACGACGTACT	TCGACTAATG	45
109 108	ORF 01	B A	ACGGCAGGCT	ATTATT	ACCT	TGCCTTGCGT	GAAATATCAA TGTA //G	CGGGGTGCGG	CAGGGGATG	52
	Methylase		CTACAACGAT CTACAAAGAT	TTTAAG	TCGG	CGCCGGGGCA		ATGTGGGGCA	GGCAACATG	104
	16S RNA	A	TCGGCGATGG CCGGCGATGG	TTTATA	TGCC	CATGGACGGG	CCGATCCGAT	CGTACGTGAC	GC.//AAT	220
ı	Archaeal p			YTTAWA			230.11000.11	JOINEGIUMC	00.77.72	

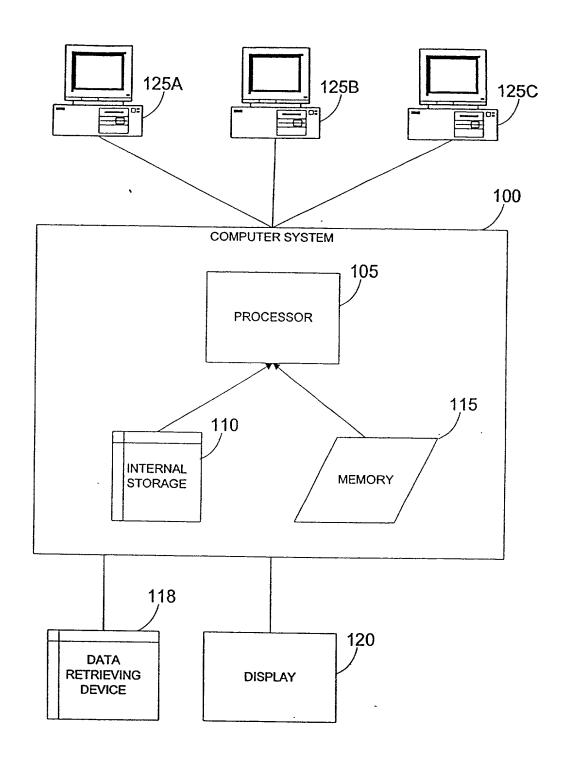


FIGURE 3

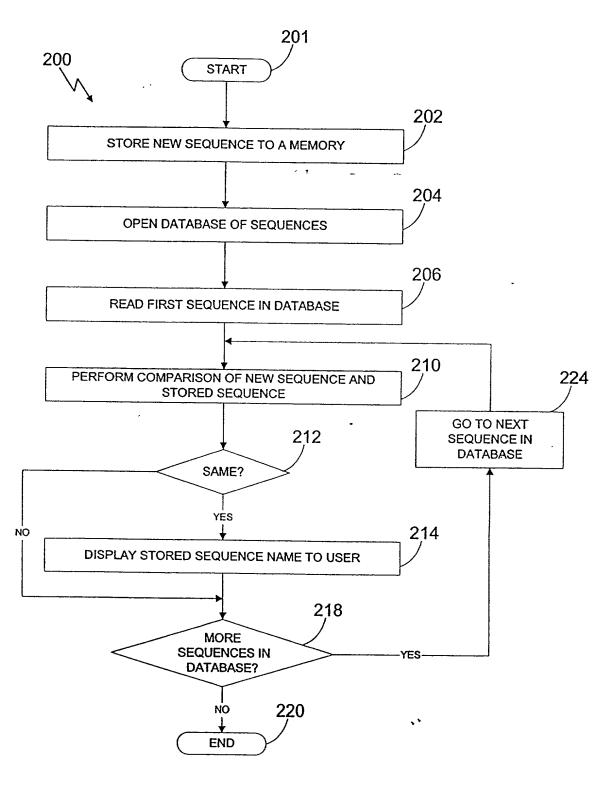


FIGURE 4

